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<110> Lal, Preeti
      Faris, Mary
      Chen, Huei-Mei
      Ison, Craig H.
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                                      40
Thr Ile Arg Leu Ile Arg Cys Gly Tyr His Val Val Ile Gly Ser
                                      55
                 50
Arg Asn Pro Lys Phe Ala Ser Glu Phe Phe Pro His Val Val Asp
                                                           75
                                      70
Val Thr His His Glu Asp Ala Leu Thr Lys Thr Asn Ile Ile Phe
                                      85
Val Ala Ile His Arg Glu His Tyr Thr Ser Leu Trp Asp Leu Arg
                 95
                                     100
His Leu Leu Val Gly Lys Ile Leu Ile Asp Val Ser Asn Asn Met
                110
                                     115
Arg Ile Asn Gln Tyr Pro Glu Ser Asn Ala Glu Tyr Leu Ala Ser
                125
                                     130
                                                          135
Leu Phe Pro Asp Ser Leu Ile Val Lys Gly Phe Asn Val Val Ser
                140
                                     145
Ala Trp Ala Leu Gln Leu Gly Pro Lys Asp Ala Ser Arg Gln Val
                155
                                     160
Tyr Ile Cys Ser Asn Asn Ile Gln Ala Arg Gln Gln Val Ile Glu
                170
                                     175
Leu Ala Arg Gln Leu Asn Phe Ile Pro Ile Asp Leu Gly Ser Leu
                185
                                     190
                                                          195
Ser Ser Ala Arg Glu Ile Glu Asn Leu Pro Leu Arg Leu Phe Thr
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                                     205
Leu Trp Arg Gly Pro Val Val Val Ala Ile Ser Leu Ala Thr Phe
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                                     220
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1

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Phe Phe Leu Tyr Ser Phe Val Arg Asp Val Ile His Pro Tyr Ala
                                     235
                                                          240
                230
Arg Asn Gln Gln Ser Asp Phe Tyr Lys Ile Pro Ile Glu Ile Val
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                                     250
                245
Asn Lys Thr Leu Pro Ile Val Ala Ile Thr Leu Leu Ser Leu Val
                260
                                     265
Tyr Leu Ala Gly Leu Leu Ala Ala Ala Tyr Gln Leu Tyr Tyr Gly
                                                          285
                                     280
                275
Thr Lys Tyr Arg Arg Phe Pro Pro Trp Leu Glu Thr Trp Leu Gln
                                                          300
                                     295
                290
Cys Arg Lys Gln Leu Gly Leu Leu Ser Phe Phe Phe Ala Met Val
                                                          315
                                     310
                305
His Val Ala Tyr Ser Leu Cys Leu Pro Met Arg Arg Ser Glu Arg
                                                          330
                                     325
                320
Tyr Leu Phe Leu Asn Met Ala Tyr Gln Gln Val His Ala Asn Ile
                                                          345
                                     340
                335
Glu Asn Ser Trp Asn Glu Glu Glu Val Trp Arg Ile Glu Met Tyr
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                                     355
                350
Ile Ser Phe Gly Ile Met Ser Leu Gly Leu Leu Ser Leu Leu Ala
                                                          375
                                     370
                 365
Val Thr Ser Ile Pro Ser Val Ser Asn Ala Leu Asn Trp Arg Glu
                                                          390
                 380
                                     385
Phe Ser Phe Ile Gln Ser Thr Leu Gly Tyr Val Ala Leu Leu Ile
                                                          405
                                      400
                 395
Ser Thr Phe His Val Leu Ile Tyr Gly Trp Lys Arg Ala Phe Glu
                                                           420
                 410
                                      415
Glu Glu Tyr Tyr Arg Phe Tyr Thr Pro Pro Asn Phe Val Leu Ala
                                                           435
                                      430
                 425
Leu Val Leu Pro Ser Ile Val Ile Leu Gly Lys Ile Ile Leu Phe
                                                           450
                                      445
                 440
Leu Pro Cys Ile Ser Arg Lys Leu Lys Arg Ile Lys Lys Gly Trp
                                      460
                                                           465
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Glu Lys Ser Gln Phe Leu Glu Glu Gly Ile Gly Gly Thr Ile Pro
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gaagtgtccg tatcatggaa tcaatctcta tgatgggaag ccctaagagc cttagtgaaa 360
cttgtttacc taatggcata aatggtatca aagatgcaag gaaggtcact gtaggtgtga 420
ttggaagtgg agattttgcc aaatccttga ccattcgact tattagatgc ggctatcatg 480
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<213> Homo sapiens

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tgagcaataa catgaggata aaccagtacc cagaatccaa tgctgaatat ttggcttcat 720
tattcccaga ttctttgatt gtcaaaggat ttaatgttgt ctcagcttgg gcacttcagt 780
taggacctaa ggatgccagc cggcaggttt atatatgcag caacaatatt caagcgcgac 840
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catcagccag agagattgaa aatttacccc tacgactctt tactctctgg agagggccag 960
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gaaagctaaa acgaattaaa aaaggctggg aaaagagcca atttctggaa gaaggtattg 1740
gaggaacaat tcctcatgtc tccccggaga gggtcacagt aatgtgatga taaatggtgt 1800
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ctgcaaggct cgccctgcc cggcgtggag ggcgcggggg gcgcggagaa agtgaagaga 180
ggaaattgga aaattgtgag tggaccttct gatactgctc ctccttgcgt ggaaaagggg 240
aaagaactgc atgcatatta ttcagcgtcc tatattcaaa ggatattctt ggtgatcttg 300
gaagtgtccg tatcatggaa tcaatctcta tgatgggaag ccctaagagc cttagtgaaa 360
cttgtttacc taatggcata aatggtatca aagatgcaag gaaggtcact gtaggtgtga 420
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taaaatcctg attgatgtga gcaataacat gaggataaac cagtacccag aatccaatgc 300
tgaatatttg gcttcattat tcccagattc tttgattgtc aaaggattta atgttgtctc 360
agcttgggca cttcagttag gacctaagga tgccagccgg caggtttata tatgcagcaa 420
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ggttattgaa cttgcccgcc agttgaattt cattcccatt gacttgggat ccttatcatc 180
agccagagag attgaaaatt tacccctacg actctttact ctctggagag ggccagtggt 240
ggtagctata agcttggcca cattttttt cctttattcc tttgtcagag atgtgattca 300
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attactttgc tctccctagt atacctcgca ggtcttctgg cagctgctta tcaactttat 180
tacggcacca agtataggag atttccacct tggttggaaa cctggttaca gtgtagaaaa 240
cagcttggat tactaagttt tatcttcgct atggtccatg ttgcctacag cctctgctta 300
ccgatgagaa ggtcagagag atatttgttt ctcaacatgg cttatcagca ggttcatgca 360
aatattgaaa actcttggaa tgaggaagaa gtttggagaa ttgaaatgta tatctccttt 420
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aatgctttaa actggagaga attcagtttt attcagtcta cacttggata tgtcgctctg 540
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PC-0037 US
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ccatccataa attaaaacat ggaaagtact tatgagcaga gcgacatatc caagtgtaga 180
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tggggaaaga gnccgatttc tggaagaagg tctgggaggg acaattcgca tgtcgccccg 180
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tgggttgaaa cttgttaaat gagatttcaa ctgacttagt gatagagttt tcttcaagtt 180
aattttcaca aatgtcatgt ttgccaatat gaatttttct agtcaacata ttattgtaat 240\,
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tgcttccgat gaccacgtga tagccgcacc tgataagccg aatggtcaga gacttggcaa 180
aatccccact tcctatcacc cccacggtga ccttccttgc gtctttgata ccgtttatgc 240
cattaggcaa aaacgtctcc agggtcttag ggcttcccat catagagatg gattccatgg 300
tagagactct tctaagatca ccaggaatgc cctgggaatc ttaaggtgta gcttctcact 360
cagaggaget ggagggagge teetteggeg etgetggaet etggaactge etacgtgtag 420
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                                                          30
Asp Thr Gly Glu Thr Ser Met Leu Lys Arg Pro Val Leu Leu His
                 35
                                      40
Leu His Gln Thr Ala His Ala Asp Glu Phe Asp Cys Pro Ser Glu
                 50
                                      55
Leu Gln His Thr Gln Glu Leu Phe Pro Gln Trp His Leu Pro Ile
                 65
                                      70
Lys Ile Ala Ala Ile Ile Ala Ser Leu Thr Phe Leu Tyr Thr Leu
                 80
                                      85
Leu Arg Glu Val Ile His Pro Leu Ala Thr Ser His Gln Gln Tyr
                 95
                                     100
Phe Tyr Lys Ile Pro Ile Leu Val Ile Asn Lys Val Leu Pro Met
                110
                                     115
                                                         120
Val Ser Ile Thr Leu Leu Ala Leu Val Tyr Leu Pro Gly Val Ile
                125
                                     130
                                                         135
Ala Ala Ile Val Gln Leu His Asn Gly Thr Lys Tyr Lys Lys Phe
                140
                                     145
                                                         150
Pro His Trp Leu Asp Lys Trp Met Leu Thr Arg Lys Gln Phe Gly
                155
                                     160
                                                         165
Leu Leu Ser Phe Phe Phe Ala Val Leu His Ala Ile Tyr Ser Leu
                170
                                     175
                                                         180
Ser Tyr Pro Met Arg Arg Ser Tyr Arg Tyr Lys Leu Leu Asn Trp
                185
                                     190
                                                         195
Ala Tyr Gln Gln Val Gln Gln Asn Lys Glu Asp Ala Trp Ile Glu
                200
                                     205
His Asp Val Trp Arg Met Glu Ile Tyr Val Ser Leu Gly Ile Val
                215
                                     220
                                                         225
Gly Leu Ala Ile Leu Ala Leu Leu Ala Val Thr Ser Ile Pro Ser
                230
                                     235
Val Ser Asp Ser Leu Thr Trp Arg Glu Phe His Tyr Ile Gln Ser
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                                     250
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525

Lys Leu Gly Ile Val Ser Leu Leu Leu Gly Thr Ile His Ala Leu

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Ile	Phe	Ala	Trp	Asn	Lys	Trp	Ile	Asp		Lys	Gln	Phe	Val	Trp
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Tyr	Thr	Pro	Pro	Thr	Phe	Met	Ile	Ala	Val	Phe	Leu	Pro	Ile	Val
				290					295					300
Val	Leu	Ile	Phe	Lvs	Ser	Tle	Len	Phe	T.011	Dro	Cys	LOU	7 200	Tara
				305	~ ~		200	1110	310	110	Cys	пец	Arg	_
T 370	τ1.	T 011	T		3	•	~-	_						315
гур	тте	ьец	гда	тте	Arg	HIS	Gly	Trp	Glu	Asp	Val	\mathtt{Thr}	Lys	Ile
				320					325				-	330
Asn	Lys	Thr	Glu	Ile	Cvs	Ser	Gln	T.011						550
	-			335	-1-	~~-	0111	Lou						
				555										